

Electronic Supplementary Data

Identification of polymorphisms in the bovine collagenous lectins and their association with infectious diseases in cattle

Immunogenetics

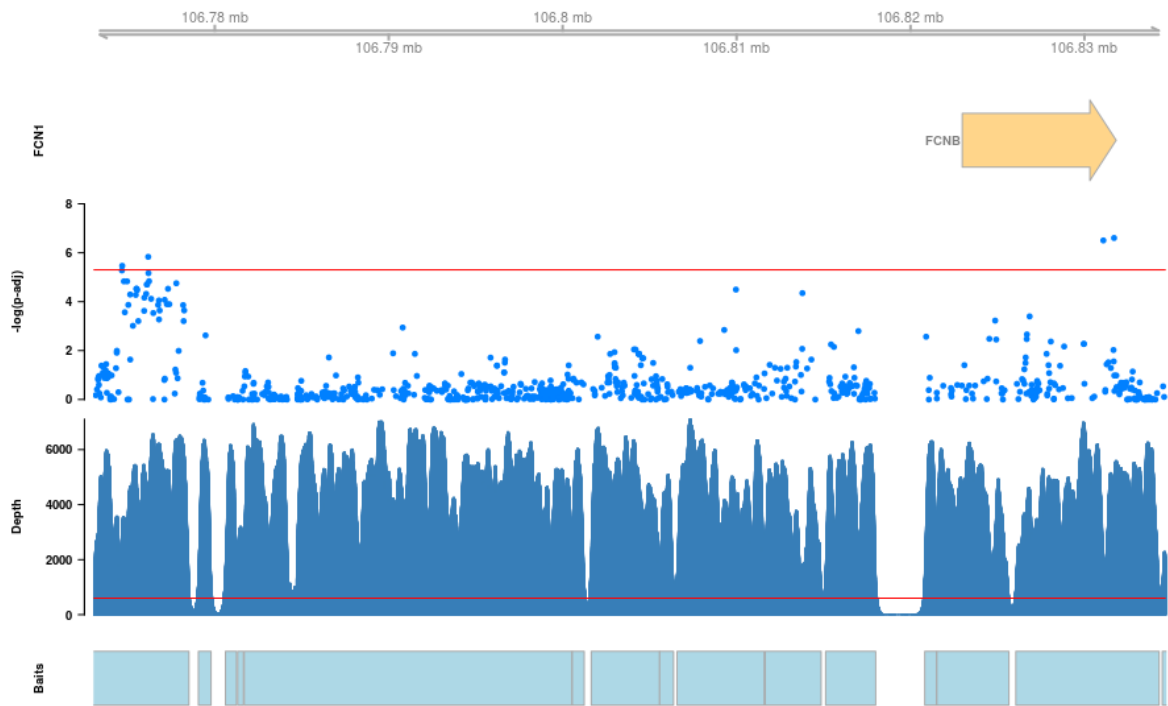
Russell S. Fraser¹, John S. Lumsden^{1,2}, Brandon N. Lillie^{1*}

¹Department of Pathobiology, Ontario Veterinary College, University of Guelph

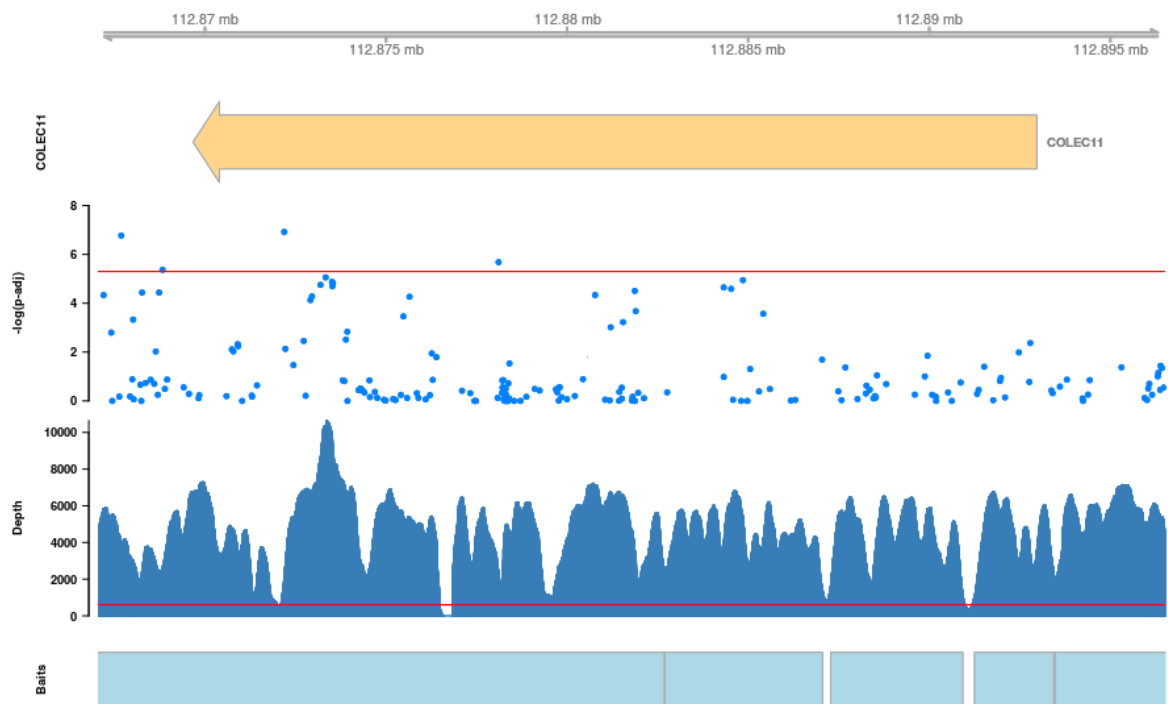
²Adjunct Professor, St. George's University, True Blue, Grenada

*Corresponding author: blillie@uoguelph.ca

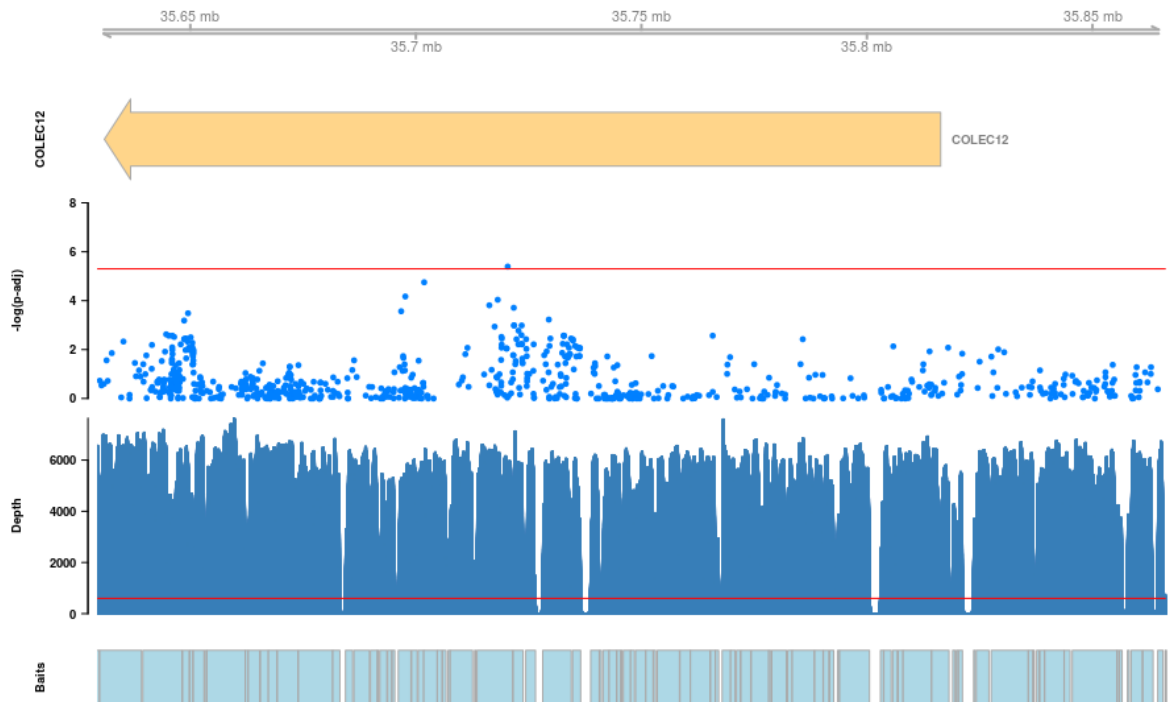
a)



b)



c)



Online Resource 2

Distribution of significant associations are shown for **a** *COLEC11*, **b** *COLEC12*, and **c** *FCN1*. The red line indicates the cutoff for BADGE class II significance ($p < 5 \times 10^{-6}$). The depth of sequencing (total from all pools) and the probes used for target capture are shown to illustrate gaps in sequencing and variant discovery.